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Bukaryota; Mctazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Mctazoa; Chordates; Catarrhini; Hominidae; Homo.

1 (bases i to 2740)

1 (bases i to 2740)

K.M., I.G., Jun. D.Y., Sohn, U. and Kim.Y.H.

Cloning and expression of human mitotic centromere-associated
                                                                                                                                               2 (bases 1 to 2740)
Kim, I.G., Jun, D.Y. and Kim, Y.H.
Direct Submission
Submitted (12-JUL-1996) Microbiology, Kyungpook National
University. Tasey 702-701, Norea
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/db_xref="taxon:9606"
/clone="E6-1"
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/product="mitotic centromere-associated kinesin"
/protein_id="AAC27660.1"
                                                                        /cell_line-"Jurkat"
                                     /note-"HSMCAK"
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ō	AACCCAAGTTGAAAGTGGACTTAACAAAG	901	DЬ
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RESULT 5 BC008764 LOCUS DEFINITION

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human, Romo sapiens Eukaryota; Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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                        CAGCGTGACTCAAACAAGAGCCAGAAACTGCTAAAGAAACTCATGTCAGGAGTGGAGAAT 860
                                                                          TCAGGCCCTTGCTTATTCTGTGGCACTCTGGTGTGTACTCATGAGGAACAAGATATTTTA 800
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Yue, H., Tang, Y. T., Hillman, J.L., Lal, P., Bandman, O., Baughn, H.R., Asilmzei, T., Yang, J., Reddy, R. and Lu, D. A. Human intra-cellular signaling molecules patent No 0077040-A 69 21-DEC-2000; Locyte Genomics, Inc. (US)
Locyte Genomics, Inc. (US)
Locyte Genomics, Inc. (US)
Locetion/Qualifiers
1. 1981
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Kammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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95.8%, Score 1913.4; DB 6; Length 1981; Similarity 98.1%, pred. No. 0; 43; Conservative 0; Mismatches 35; Indels 3; Gaps <u>ب</u>

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ICAATTGAGAGTGCTGAAGAGATACGAGAATATGTTACTGATCTCCTCCA-GGGAAAT 179

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proc. Natl. Acad. Sci. U.S.A.
86094327
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PD 12-AU
PP 31-JA,
PI KARAG
PC C12R5:91
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Adwayuchi, R. and Hiklchi, K. CELL-LINE ORIGINATED FROM HUMAN BONE MARROW CELLS Patent: JP 1997206070-A 1 12-AUG-1997;
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tion replace(1317,'a'),
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                                                                                                                                                                                                                                      Contact:
Shevcherko, Y. Wetherby, E.D. Beckstom Sternberg, S.M.,
Beolman, B. Blakesley, R. W. Bourkston, Brinkley, C., Brooks, S.,
Dietrich, H. D., Guan, X., Guyta, J. Hard, S. B. Hardton, E. Contagall, R.,
Kinn, H., Haduro, O.L., Massello, C., Mastrin, E. D., McCloskey, J.C.,
KCDowell, J., Pearson, R. Sydder B., Stantipop, S., Thomas, J.,
Tiongson, E.E., Touchman, J.W., Tsurgeon, C., Vgy, J.L., Malker, M.A.,
                                                                                                                 through the I.H.A.G.E. Consortium/LLML at: http://lmage.llnl.gov
Series: IRAL Plate: 12 Row: m Column: 12
This clone was selected for full length sequencing because it
passed the following selection criteria: matched mRNA gl: 4505190
                                                                                                                                                                                                                                                                                                                                                                                               CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: National Institutes of Health Intramural
Sequencing Center (NISC),
                                                                                                                                                                                         Clone distribution: MGC clone distribution
                                                                                                                                                                                                                                                                                                                                                               Gaithersburg, Maryland;
Web site: http://www.nisc.nih.gov/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA

NIH-MGC Project URL: http://mgc.nci.nih.gov

Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
Submitted (05:FEB-2001) National Institutes of Health, Mammalian
Gene Collection (MCC), Cancer Genomics Office, National Cancer
                                                                                                                                                                                                                                 Zhang, L.-H. and Green, E.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-remail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1. (Dases 1 to 2745)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Institute, 31 Center Drive, Room 11A03, Bethesda,
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Organism-"Romo santems"
/th_xref-"taxn.9600"
/clone-"MG:3690 IMMGE:3629765*
/tissue_type-"Placenta, Chorlocarcinoma"
/clone_lib-"NH_MGC_216
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Run on:
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July 27, 2002, 05:34:43; Search time 3956.53 Seconds (Without allgaments) 1351.930 Million cell updates/sec
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Title: Perfect score: Sequence: PCT-US02-13994-3 2885 1 ggmattcctcttgtc gyaattootottgtogaagt.....aaacaaaagttaaaaattt 2885

Scoring table: 1736436 seqs, 858457221 residues IDENTITY_NUC Gapop 10.0 , Gapext 1.0

Total number of hits satisfying chosen parameters: 3472872

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

N. Geneseq. 032802:

N. SIDSI 9990data/no
SIDSI 9990data/no 1. SIDSI/Gogdata/hold-geneseq/geneseqn-embl/NA1981.DAT:
3. SIDSI/Gogdata/hold-geneseq/geneseqn-embl/NA1981.DAT:
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8. SIDSI/Gogdata/hold-geneseqn-genesqn-embl/NA1991.DAT:
8. SIDSI/Gogdata/hold-genesqn-genesqn-

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

987552	Result
2885 2831.4 2181.6 1558.4 953.6 6671 669.6 668 667	Score
100.0 98.1 75.6 54.0 33.1 23.3 23.2 23.2 23.2 23.2	Query re Match Length DB
2885 4021 2233 2601 4189 4189 3208	Length
222222222	뛿
AAX40055 AAC77217 AAC8958 AAX89456 AAX894456 AAX59647 AAC77054 AAX51858 AAX18086	ID
Colon cancer assoc Human DEFX DEF2772 Human histone deac Human Lill-length Human DEFX DEF269 Human CDNA sequenc	Description

	AAH06343	22	638	ω •	97.6	4.5
Human breast cance	AAL15037	22	493	.2	121.4	1
Drosophila melanog	ABL10648	23	9384	4.7	134.6	4
Human immune/haema	AAK57190	22	325	4.7	135.2	12
Arabidopsis thalia	AAC50614	21	1993	. 8	138	-
Arabidopsis thelia	AAC34680	21	1991	. 8	138	40
	AAL23890	22	501	B	138.6	39
Human immune/haema	AAK69720	22	323	B	139.2	38
cDNA encoding nove	AAS41213	22	606	5.2	148.6	37
Human CDNA sequence	AAH18350	22	2552	٠ <u>.</u>	151.8	36
cDNA encoding nove	AAS41280	22	1005	5.3	152	35
Human ORFX ORF2507	AAC76952	21	2480	5.4	155.2	u A
Human ORFX ORF2521	AAC76966	21	2461	5.	156.8	33
Human musculoskele	AAL36019	22	5930	5.5	159.8	32
Human ORFX ORF3078	AAC77523	21	995	5.6	161.8	2
Drosophila melanog	ABL10649	23	4689	5.6	163	30
	AAK60936	22	359	6.0	173	29
Human DNA sequence	AAS95017	2	6949	6.0	173.6	87
Human cDNA sequenc	AAH14416	22	2548	6.1	175.2	27
Murine transcripti	AAK53847	22	271	6.1	177.2	26
	ABL03352	23	22341	7.4	214.6	25
	AAH18707	22	5020	7.5	216.8	24
	AAK69721	22	16086	10.0	289	23
TUSC!	AAL36020	22	16086	10.0	289	22
CDNA	AAH07464	22	719	12.1	350.2	21
Human Char Sequence	AAH03697	22	606	12.9	373.2	20
200	AAH16517	2	1880	13 3	383.6	19
	AAH07105	2	607	13.6	391.4	18
con encoung nove	ABI.03353	2	3894	13	401.2	17
manda con sequence	AAS41232	2	1085	14.6	421.8	16
	AAH13829	22	2026	15.7	452.6	15
2	AAH16356	22	1745	19.4	60.	1
	AAK93407	22	626	19.7	6	13
Himan CDNA 5' and	AAK92144	22	626	19.7	33	12
_	AAC81740	21	2621	20.0	77	11
Human bolynucleott	AAK53008	22	4200	22.8	558.6	10

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ALIGNMENTS

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RESULT	LT 1				
AAX40055	0055				
D	AAX40055 standard; DNA;	DNA:	2885	BP.	
χÃ	AAX40055;				

Colon cancer associated gene. 02-JUL-1999 (first entry)

Cancer associated antigen; diagnosis; research; treatment; human; breast cancer; colon cancer; gastric cancer; renal cancer; lung cancer; prostate cancer; ss.

22-JUN-1998; 17-JUL-1997; 10-OCT-1997; 10-OCT-1997; 10-OCT-1997; 11-OCT-1997; (LUDW-) LUDWIG INST CANCER RES 28-JAN-1999. W09904265-A2 Homo sapiens 15-JUL-1998; 98US-0102322. 97US-0896164. 97US-0061599. 97US-0061765. 97US-0948705. 97GB-0021697. 98WO-US14679.

Chen Y, Gout I, Gure A, O'Hare M, Obata Y, Old L

Pfreundschuh M. Sahin U. Scanlan MJ. Stockert E: Tureci O;

WPI; 1999-132448/11.

New isolated cancer associated nucleic acids and polypeptides - isolated using sera from cancer patients, used to develop products for the diagnosis, monitoring or treatment of cancers

Claim 67; Page 651-652; 787pp; English.

THE KKE SERVICE OF SOUTH OF THE KKE SERVICE OF THE SERVICE OF THE KKE SERVICE OF THE KKE SERVICE OF THE SERVI

The invention relates to a method for diagnosing a disorder characterised by expression of a human cancer associated antiqen precursor coded for by a nucleic acid molecule (NAM). The method comprises: (a) contacting a binds to the ANM, an expression product or a fragment of an expression product complexed with an HLA molecule; and (b) determining the interaction between the agent and the NAM or the expression product as a determination of the disorder. The products and methods can be used in the diagnosis, monitoring, research, or treatment of conditions the disorder, monitoring or various cancer associated antiqens, which are cancer associated antigen precursors expressed in human breast cancer, renal cancer, colon cancer, gastric cancer, prostate cancer and lang cancer.

Sequence 2885 BP; 626 A; 901 C; 838 G; 520 T; 0 other;

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gaacgagocticoctotygacagotcoccaaccagttcagcotctacacgtctcottc geacgagocctcoctctygacagotcoccaaccagttcagcotctacacgtctcottc geacgagocctcoctctygacagctcocccaaccagttcagcotctacacgtctcottc
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Score 2885; Pred. No. 0; Mismatches

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31-MAR-1999; 02-APR-1999; 05-APR-1999; 30-MAR-2000; 2000US-0540763 99US-0127636. 99US-0127636. 99US-0127728.

31-MAR-2000; 2000WO-US08621

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> Shimkets RA, (CURA-) CURAGEN CORP Leach M;

P-PSDB; 2000-602362/57 AAB43008

Novel nucleic solds and peptides derived from open reading useful for treating e.g. cancers, proliferative disorders, neurodegenerative disorders and cardiovascular disease.

Claim 5; Page 4726-4728; 5507pp; English

AAC74446 to AAC77606 encode the protein given in Abs40337 to Abs413397. Which represent the human ORFX open reading frames it to 3161. The 3097 sequences have activaties such as cytostactic to 2161. The 3097 enripportatic antipartition and incomplete such as cytostactic unexponentive: otherwise antipartition and incomplete such as cytostactic unexponentive: osteopathic; anticomplete such antipartition of the sequence of the sequence of the sequence of the sequence of the sequences on be used for determining the presence of or predisposation of the sequences on be used for determining pathological conditions associated with an ORFX-associated disorder. The

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Pfreundschuh M, Sahin ď, Scanlan MJ, Stockert E,

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for the diagnosis, monitoring or treatment of cancers New isolated cancer associated nucleic acids and polypeptides solated using sera from cancer patients, used to develop products

Claim 67; Page 677-678; 787pp; English.

The Invention relates to a method for diagnosing a disorder characterised by expression of a human cancer associated entiren precursor coded for by a nucleic acid molecula (NAM). The method companies: retocating a hological sample isolated from a retocating a hological sample isolated from a roduce at fragment of an expression binds to the NAM; an expression product complexed with an entire roduce to the second product as a network of the second product and network of the second product as a network of the second product as a network of the second product as a network of the second product and network of the second product as a network of the second product and network of the second cancer.

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Homo sapiens

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           Human; chromosome mapping; gene mapping; gene therapy; forensic;
food supplement; medical imaging; diagnostic; genetic disorder;
                                                           DNA encoding novel human diagnostic protein #23250
                                                                                             13-FEB-2002
                                                                                                                                                           AAS87446 standard;
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diagnostics, forensics, gene mapping, identification of mutations
responsible for genetic disorders or other traits and to assess
biodiversity.
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23-AUG-2000; 2000US-0649167
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Sequence 2265 BP; 784 A; 356 C; 491 G; 634 T; 0 other;

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OM nucleic - nucleic search, using sw model
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Run 9 July 27, 2002, 05:39:00; Search time 3956.53 Seconds
471.698 Million cell updates/sec

a

Title: Perfect score: Sequence: PCT-US02-13994-B 1087 1 aagatgatgcctag aagatgatgcctagtaaatt.....aataaatgattatattcaca 1087

Scoring table: IDENTITY_NUC Gapop 10.0 , Gapext 1.0

Total number of hits satisfying chosen parameters:

3472872

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1736436 seqs, 858457221 residues

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Listing first 45 summaries Maximum Match 100%

Database

N. Geneseq_03203.*

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ALIGNMENTS

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standard;		

DNA; 1087 BP.

AAX40070;

02-JUL-1999 (first entry)

Colon cancer associated gene.

prostate cancer; ss. Cancer associated antigen; diagnosis; research; treatment; human; breast cancer; colon cancer; gastric cancer; renal cancer; lung cancer;

Homo sapiens

W09904265-A2.

28-JAN-1999.

22-JUN-1998; 17-JUL-1997; 10-OCT-1997; 10-OCT-1997; 10-OCT-1997; 11-OCT-1997; 15-JUL-1998; 98US-0102322. 97US-0896164. 97US-0061599. 97US-0061765. 97US-0948705. 97GB-0021697. 98WO-US14679

(LUDW-) LUDWIG INST CANCER RES

Chen Y, Gout I, Gure A, O'Hare M, Obata Y, PTO PTO

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News isolated cancer associated nucleic acids and polypeptides -isolated using sera from cancer patients, used to develop products for the diagnosis, monitoring or treatment of cancers

Claim 67; Page 676-677; 787pp; English.

blinds to the NAM, an expression product or a fragment of an expression product complexed with an HLA molecule; and (b) determining the interaction between the agent and the NAM or the expression product as a determination of the disorder. The products and methods can be used the disorders are products and methods can be used in the disorders are products and methods can be used in the disorders are producted and the products are all the disorders and produced and the disorders are characterised by the expression of various cancer associated antiques which are cancer associated antique produces and encoded polypeptides which are cancer associated antique products are producted cancer. Journal of the concert of the co The invention relates to a method for diagnosing a disorder characterised by expression of a human cancer associated antigen precursor coded for by a nucleic acid molecule (NAM). The method comprises: (a) contacting a biological ample isolated from a subject with an agent that specifically blids to the NAM, an expression product or a fragment of an expression

Sequence 1087 BP; 386 ? 212 C; 203 <u>و</u> 286 T; 0 other;

100.0%;

Score 1087; Pred. No. 1.

1.2e-265;

DB 20; Length 1087;

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ID AAV AAV59766 standard; DNA; 1455 BP

AAV59766;

(first entry)

Human secreted protein gene 106 clone HT3AM65

cognitive disorder; schlsophrenia; prostate; obesity; osteoclast; thymus osteoporosia: arthuttis; testis; lung; thyroid/tils; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm; Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative olisorder; butwania; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthas; lymphocytic disease; brain; hepatic; lymphoma; infilammation; ischaemic shock; Alzheiner's disease; restenosis; AlDS;

sapiens

W09839448-A2

11-SEP-1998

06-MAR-1998; 98WO-US04493

02-OCT-1997 07-MAR-1997 07-MAR-1997 07-MAR-1997 07-MAR-1997 07-MAR-1997 07-MAR-1997 07-MAR-1997 07-MAR-1997 07-MAR-1997 970S-0061060. 970S-0038621. 970S-0040161. 970S-0040162. 970S-0040163. 970S-0040333. 970S-0040336. 970S-0040336.

GenCore version 4.5 Copyright (c) 1993 - 2000 Comp Compugen Ltd

OM nucleic nucleic search, using sw model

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July 27, 2002, 05:40:04; Search time 3955.53 Seconds
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Title: Perfect score: Sequence: PCT~US02-13994-10 1953

Scoring table: IDENTITY_NUC Gapext 1.0 acgcctgccaggagcaagcc.....aaagaatgttttggaacttt 1953

1736436 seqs, 858457221 residues

Total number of hits satisfying chosen parameters:

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

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SUMMARIES

Fred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

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1953 1791.2 1253.2 1253.2 577.6 544.8 482.4	Score
100.0 100.0 91.7 64.2 29.6 29.6 27.9 24.7 24.7	Query Match
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AAX39673 standard; DNA; 1953 BP.

AAX39673;

02-JUL-1999 (first entry)

Renal cancer associated gene.

Cancer associated antigen; diagnosis; research; treatment; breast cancer; colon cancer; gastric cancer; renal cancer; human; lung cancer,

prostate cancer; ss.

Homo sapiens

W09904265-A2 28-JAN-1999

15-JUL-1998; 98WO-US14679

22-JUN-1998; 17-JUL-1997; 10-OCT-1997; 10-OCT-1997; 10-OCT-1997; 11-OCT-1997; 980S-0102322. 970S-0896164. 970S-0061599. 970S-0061765. 970S-0948705. 970S-0021697.

(LUDW-) LUDWIG INST CANCER RES

Chen Y, Gout I, Gure A, O'Hare M, Obata Y, 01d LJ

Pfreundschuh M, Sahin U, Tureci O; Scanlan MJ, Stockert E;

New isolated cancer associated nucleic acids and polypeptides -isolated using sera from cancer patients, used to develop products for the diagnosis, monitoring or treatment of cancers diagnosis, monitoring or treatment of cancers

REXERRE EXERTING SOCIOSOS SOCIOS SOCI Claim 67; Page 485; 787pp; English.

The invention felates to a method for diagnosing a disorder characterised by expression of a human cancer associated antique precursor coded for by a nucleic acid molecule (NMM). The method comprises rely contexting a nucleic standard form a subject to the same of the specifically binds to the NAM, an expression product or a fragment of an expression product complexed with an HAM more than the decreasion product to a fragment of the discount of the discount and the NAM or the expression product as a determination of the discount or products and methods which are cancer associated actigen precursors expressed in human breast mention provides nucleic acid sequences and encoded polypoptions which are cancer associated antigen precursors expressed in human breast maner. Family cancer, renal cancer, colon cancer, gastit cancer, prostate cancer and lung cancer.

Sequence 1953 BP; 429 A; 580 C; 654 G; 290 T; 0 other;

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Mon Jul 29 12:11:49 2002

GenCore version 4.5 Copyright (c) 1993 - 2000 Comp Compugen Ltd

OM nucleic - nucleic search, using sw model

Run

on:

July 27, 2002, 05:48:21; Search time 3956.53 Seconds
Without alignments
1824.303 Millon cell updates/sec

Title: Perfect score:

Scoring table: Sequence: PCT-US02-13994-14
4204
1 acycaggcagtgatgtcacc.....gagtagtggagtgtcagtgc 4204

IDENTITY_NUC Gapop 10.0 , Gapext 1.0

Searched: 1736436 seqs, 858457221 residues

Total number of hits satisfying chosen parameters: 3472872

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

N. Genesse, 032802:*

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Pred. Mo. is the number of results predicted by chance to have a score of the result being printed and is derived by analysis of the total score distribution.

SUMMARIES

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Tumour rejection a	Human MAGE-12 gene	MAGE-2 gene. Homo	MAGE-2 gene. Homo	Tumour rejection a	DNA encoding molec	Human melanoma ass	Human MAGE-A3 nucl	cDNA encoding MAGE	Description	

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AAX26974 standard; cDNA; 4204 BP.

AAX26974;

25-JUN-1999 (first entry)

cDNA encoding MAGE-3 polypeptide.

MAGE-3 tumour associated gens; human leucocyte antigen Class II; autologous CD4+ coli, MAGE-3 related disease; cancer; melanoma; osteosarcoma; leukemia; carcinoma; ss.

Homo sapiens

CDS Location/Qualifiers 2465..3409 /product= "MAGE-3"

W09914326-A1

25-MAR-1999.

04-SEP-1998; 98WO-US18601

12-SEP-1997; 9705-0928615

(LUDW-) LUDWIG INST CANCER RES (UYVR-) UNIV VRIJE BRUSSEL.

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Boon-Falleur T, Chaux P, Corbhals J,
Luiten R, Stroobant V, Thielemans K,
P-PSDB; AAY01720.
                     WPI; 1999-244031/20.
                                                                Heirman C;
Van Der Bruggen P;
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Example 5; Page 65-67; 88pp; English.

molecules

Isolated peptides that bind to human leucocyte antigen class II

The present sequence represents the MAGE-3 tumour associated genepertides that bind human leucocyte antigen (HLA) Class II molecules
can be derived from the MAGE-3 protein. These peptides and
and HLA Class II. are used to treat MAGE-3 peptide
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particularly cancers (e.9. melanoms, osteosarcoms, leukemia and
particularly cancers (e.9. melanoms osteosarcoms, leukemia and
particularly bettection of of the peptides, e.g. in binding
assays, particularly with antibodies, is used for diagnosis of such diseases.

Sequence 4204 BP; 944 A; 1144 C; 1223 G; 893 T; 0 other;

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                 The present invention relates to MAGE-Al (tumour associated gene product) human leukoyte antigen (HA) class II-binding peptides (see AB80256-80,2595, and AB8025631-902637). These peptides are presented to cells in the context of HA class III molecules. The peptides stimulate the activity and proliferation of CD4-T lymphoytes. The invention also includes nuclectide sequences encoding MaGE-3 peptides (see AAA7938 and AAA7938-A79940). The peptides and nuclectide sequences can be used to create antibodies against the MAGE-3 peptides, the antibodies. The peptides are used to dispose or treat a disorder characterized by eptides are used to dispose or treat a disorder characterized by apprides are used to dispose or treat a disorder characterized by apprides are used to dispose or treat a disorder characterized by apprides are used to dispose or treat a disorder characterized by apprinces of the methods can also be used appreciated to the contract of 
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tumours, are fragments of MAGE-A3 which bind to and are presented to
lymphocytes by human leukocyte antigen class II molecules
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Bakaryota, Medazoa, Chordata, Craniata, Vertebrata, Exteleostomi, Mammalia; Extheria, primates, Catarrhini; Mominidae; Homo. 1 (bases i to 766) [Fisher,C., Gill,S., Grimer,R., Chand,A., Shipley,J., Gasterson,B.A. and Cooper,C.S.
                                                                         Homo sapiens
                                                                                                    SSX1 gene; SSX2 gene; synovial sarcoma
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Direct Submission
Direct Submitted (10-ANR-1995) C.S. Cooper, Institute of Cancer Research,
Submitted (10-ANR-1995) C.S. Cooper, Institute of Cancer Research,
Haddow Laboratories, Cotswold Road, Sutton, Surrey SM2 SM2, UK
Haddow Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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/ FOODO, STATT
/ FOODOATARPTYGAGIPEKIGNAFDDIAKYFSKEEWEKHKASE
KIFYYMRKKTEACHTKLOFKALDFPKCHKRAEDFGGNULDNDNRGNUTERGOFTFG
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Pfreundschuh, M.
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Patent: US 5798264-A 2 25-AUG-1998;
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